Novel serine proteinase inhibitor of the Kunitz family, BTL.010 useful for treating emphysema, cystic fibrosis, adult respiratory distress syndrome, rheumatoid arthritis, organ failure and glomerulonephritis.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC acid sequence being a generic sequence for a Kunitz-type protease
CC inhibitor or a variant where the sequence is at least 80% identical to
CC the kunitz domain of human kunitz type protease inhibitor HKI-18. Also
CC included are an isolated polypeptide obtainable by cultivation of a host
CC included are an isolated polypeptide obtainable by cultivation of a host
CC included are an isolated polypeptide construct encoding kunitz-type protein
CC in an appropriate growth medium under conditions allowing expression of
CC culture medium, a polynucleotide construct encoding the kunitz-type protein
CC culture medium, a polynucleotide construct encoding the kunitz-type
CC culture medium, a polynucleotide construct encoding the kunitz-type
CC culture medium, a polynucleotide construct encoding the kunitz-type
CC culture medium, a polynucleotide construct encoding the kunitz-type
CC culture medium, a polynucleotide construct. The
CC shock syndrome, dissedinated intravascular coagulation, hyperfibrinolytic
CC shock syndrome, dissedinated intravascular coagulation, hyperfibrinolytic
CC shock syndrome, cardiopulmonary bypass (CPB)-induced pulmonary injury,
CC major surgery, cardiopulmonary bypass (CPB)-induced pulmonary injury,
CC allergy-induced protease kelease, deep vein thrombosis, emphysema,
CC inflammatory bowel disease, and psoriasis. The present sequence
CC culture beautiful to a polynucleotide construct.
CC culture medium, a polynu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human BTL.010; neutral serine protease inhibitor; elastase inhibitor; proteinase-3 inhibitor; Kunitz domain; emphysema; idiopathic pulmonary fibrosis; adult respiratory distress syndrome; cystic fibrosis; rheumatoid arthritis; organ failure; glomerulonephritis; platelet activation; blood coagulation; neutrophil activation; neutrophil activation; neutry;
                   WPI; 2001-190860/19
N-PSDB; AAF59750.
                                                                                      Davies C,
                                                                                                                                                                                                                                      05-AUG-1999;
                                                                                                                                                                                                                                                                                   05-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                 US6180607-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                            (DAVI/) DAVIES C. (CHEN/) CHEN D.
                                                                                                                                                                                                                                                                                                                                  30-JAN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vascular injury;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human protease inhibitor BTL.010 Kunitz domain, SEQ ID NO:1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB60623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB60623 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Fig 8; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disprders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5<u>4</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    52;
                                                                                                                                        ROCZNIAK S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 YPVRCTLPPATGSCKAWAARWYFVASVGQCNRFWYGGCHGNANNPASEQECMSSCQGS 111
                                                                                    Chen D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                   99US-00369494
                                                                                                                                                                                                                                                                                   99US-00369494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       nephrotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 58
                                                                                           Roczniak S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88.4%;
89.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 391; DB 6;
Pred. No. 7.2e-27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antirheumatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
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ص
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pulmonary fibrosis, adult respiratory distress syndrome, cystic fibrosis, rheumatoid arthritis, organ failure or glomerulonephritis. BTL.010 compositions of the invention modulate at least one physiological condition such as platelet activation, blood coagulation, neutrophil activation, or monocyte activation. BTL.010 is also useful for the prophylactic or therapeutic treatment of patients undergoing angioplasty, and for the treatment of inflammatory diseases and diseases involving lung and vascular injury. The present sequence represents the human BTL.010 protease inhibitor Kunitz domain
                                                                                                                                                                                                                                                                                          database using the Kunitz domain sequences AAB60630, and was confirmed as being novel using the Kunitz domain sequences AAB60632, and AAB60633. This sequence information was extended to provide a larger region of BTL.010 protein sequence data (AAB60634) by identifying an open reading frame (ORP) which comprised DNA encoding the BTL.010 Kunitz domain fragment in a 399 bp fragment of human genomic DNA (AAF59750), corresponding to bases 16016-16414 of GenBank accession number AC004846. The entire BTL.010 Kunitz domain sequence (AAB60623) was obtained from the BTL.010 For encoded sequence. The BTL.010 protein, and pharmaceutical compositions comprising it, may be used for inhibiting protease activity, particularly that of leukocyte elastase, in the prevention, treatment or amelioration of medical conditions such as emphysema, idiopathic
Sequence 58 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               trypsin-like and chymotrypisn-like proteases. A substantial proportion o the BTL.010 protein Kunitz domain (AAB60631) was identified via homology searching in the GenBank high throughput genomic (HTG) DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 neutral serine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB60634).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 6; Col 9-10; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kunitz family,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ily, BTL.010 (fragments given in AAB60623, AAB60631 and The BTL.010 protein is thought to preferentially inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        proteases such as elastase and proteinase-3, relative to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a novel human serine protease inhibitor of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of.
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밁 Matches Local Similarity 48; 1 YPVRCLLPPATGPCKARIIRWYFVASVGQCNRFVYGGCRGNANNFASEQECMSSCQGS 58 YPVRCLLPSAHGSCADWAARWYFVASVGQCNRFWYGGCHGNANNFASEQECMSSCQGS 58 Conservative 80.2%; ° Score 264; DB 4; Pred. No. 6.3e-24; Mismatches 10; Indels 0 Gaps 0

Length 58;

Query Match

BG71912 standard; protein; 58

Human Kunk 22-JAN 2003 tz protease inhibitor protein HKI-18 (first entry)

Human; protease inhibitor; kunitz; HKI-18; antiinflammatory; anticoagulant; coagulant; CBP; psoriasis; emphysema, systemic inflammatory response syntone; acute pancreatitis; about coagulant. Homo sapiens. cardiopulmonary bypaes-induced pulmonary injury; rheumatoid arthritis; allergy-induced protease release; deep vein thrombosis; shock syndrome; & hyperfibrinolytic respiratory distr Nisseminated intravascular coagulation; ic baemorrhage; myocardial infarction; 5. .55 /label= Kunitz_domain /note= "This domain" location/Qualitiers syndrome; chronic claimed in inflammatory bowel disease claim 18"

WO200296938-A2